

GEOGRAPHICALLY DISTINCT BUT NON-MONOPHYLETIC MORPHS: REEXAMINING
THE EVOLUTION OF COLOR IN FLORIDA BURROWING BEETLES (COLEOPTERA:
GEOTRUPIDAE: *PELTOTRUPES*)

By

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**Geographically distinct but non-monophyletic morphs: Reexamining the evolution of color
in Florida burrowing beetles (Coleoptera: Geotrupidae: *Peltotrupes*)**

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ABSTRACT - The genus *Peltotrupes* Blanchard 1888 (Coleoptera: Geotrupidae) is restricted to scrub and sandhill habitats on peninsular Florida's ridge system. Commonly called Florida Deep Digger Scarabs, individuals excavate burrows as deep as 2-3 m and were traditionally considered to be flightless. The recognized diversity within the genus (*P. profundus profundus* Howden 1952, *P. p. dubius* Howden 1955, and *P. youngi* Howden 1955) is based mostly on structural color differences corresponding to distinct geographic regions (i.e., dune ridges). Herein, we provide the first molecular phylogeny of the genus *Peltotrupes* and highlight the lack of monophyly of the currently recognized species (i.e., color morphs). We also analyze the color of four dorsal regions of male individuals representing the full geographic and color phenotypic breadth of the genus. Although significant color differences are recovered between the currently recognized species, given their lack of monophyly, our results suggest an ecological role may be maintaining these distinct morphs.

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Introduction

Background: The beetle genus *Peltotrupes* (Blanchard, 1888) is native to peninsular Florida and is composed of two species, *P. profundus*, and *P. youngi* (Howden, 1952). They are seemingly flightless beetles and are known for their iridescent coloration. Additionally, they are known as deep digger scarab, and dig deep burrows to rear their young and inhabit during the day time. Along with these species, there is a subspecies, *P.p dubius* (Howden, 1955). Although *P.p. dubius* subspecies has been classified as a subspecies of *P. profundus*, this was originally dismissed and has been questioned since. These species were once classified under the genus *Geotrupes*, but this was changed to *Peltotrupes* by Henry Howden after he divided the genus into the two species: *profundus* and *youngi* (Howden, 1952). *Peltotrupes profundus* generally have a purple coloration along their elytra while *P. youngi* are typically green. The independent species within *Peltotrupes* with their robust, iridescent-coloration occupy separate geographic regions within central Florida (Kalisz, et al., 1984). Additionally, degree of coloration and other anatomical structures make these beetles sexually dimorphic (Howden, 1955).



Figure 1. *P. youngi* in sandhill habitat.



Figure 2. *P. p. profundus* in sandhill habitat.

These beetles are considered to be distinct species; however, through recent molecular phylogenetic analysis (Brewer lab) indicates the two species are not reciprocally monophyletic. These new data raise new questions regarding their physical characteristics that distinguish the species. Given the inconsistencies between genetic and morphological data, the purpose of this study was to examine color variation within and between the two species.



Figure 3. *Peltotrupes* mound habitat.



Figure 4. Up-close mound and burrow entrance.

Research Questions: (1) Do the nominal *Peltotrupes* taxa comprise reciprocally monophyletic groups? (2) Does the minor coloration used to delineate species and subspecies significantly differ?

Methods

Collection: Beetle specimens were collected from sandhill regions throughout peninsular Florida, and specimen names, species, and localities were recorded in Excel (Table 1). These beetles were collected in pitfall traps (buried coffee cans baited with sucrose/yeast mixture).

Molecular Work: Beetles collected were euthanized and pinned, and the right rear leg was removed for DNA extraction using Qiagen's DNeasy kit. After extraction, the cytochrome oxidase 1 (COX1) and cytochrome oxidase 2 (COX2) genes were amplified using Polymerase Chain Reaction (PCR). After additional clean up and gel electrophoresis, the samples were sequenced, and a SPLITSTREE phylogenetic network of mitochondrial haplotypes (Figure 8).

Photography: We generated composite photographic images for 101 male *Peltotrupes* beetle specimens representing the wide-range distribution of the genus for color analysis. Specimens were photographed using a Canon 6D camera under controlled light settings to minimize glare and ensure all specimens had similar lighting. Eight to twelve photographs were taken of each specimen and were exported to Helicon Focus, which pieced all images together to create one high definition composite image used for color analysis.



Figure 5. High-definition photograph of *Peltotrupes youngi*. Image used for color analysis.



Figure 6. High-definition photograph of *Peltotrupes p. profundus*. Image used for color analysis.



Figure 7. High-definition photograph of *Peltotrupes p. dubius*. Image used for color analysis.

Color Analyses: Composite images were exported to GIMP, a photograph editing software. The GIMP “color-picker” tool was used to measure RGB values for four dorsal areas of each specimen and a standardized grey background. The four dorsal areas included: lateral elytrum, edge elytrum, anterior, and head (Figure 8). RGB values standardized by dividing each value by the corresponding grey background value, then multiplied by 255, the RGB value of white (The protocol of measuring color values is described in the Appendix). RGB values were then converted to hexadecimal values and integer values using the R packages RGB and STROI to provide single

continuous values for analysis. Specimens, grouped by taxon, were subjected to ANOVA, followed by Tukey's post-hoc tests in R. Lastly, principal component analysis (PCA) was used to cluster taxa using all four dorsal measurements and subjected PERMANOVA to test for differences between taxa (999 iterations).

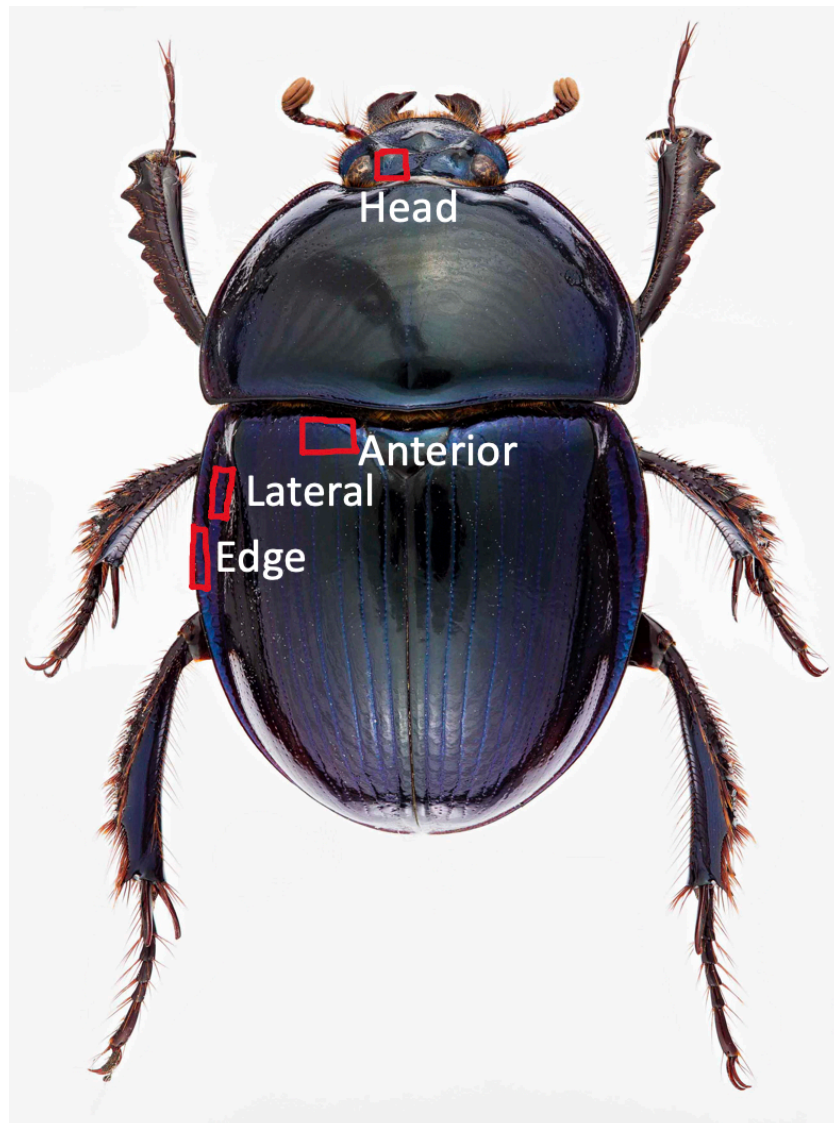


Figure 8. *Peltotrupes profundus* illustrating locations measured for color.

Results

Brief Synopsis of Molecular Work: The SPLITSTREE phylogenetic network revealed phylogeographic structuring among mtDNA haplotypes, with four distinct phylogroups (Figure 9). However, none of the three nominal taxa were resolved as being monophyletic. Instead, all of the haplotypes representing *P. youngi* and *P.p. dubius* clustered with those of *P. p. profundus* from the northern portion of its geographic range .

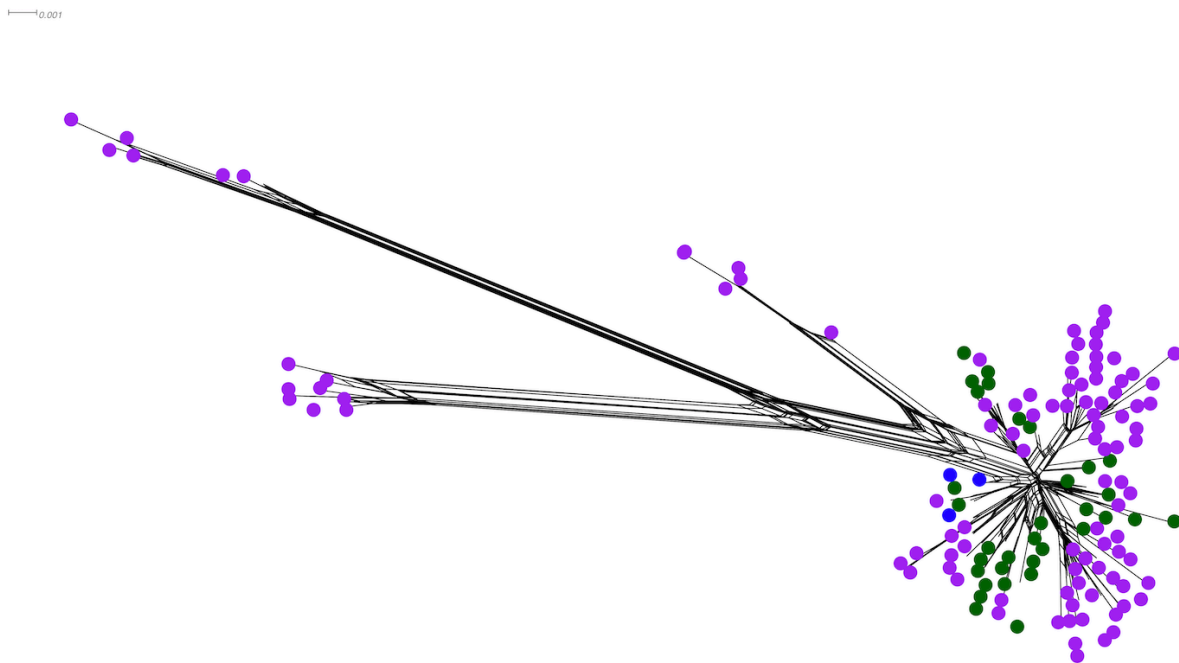


Figure 9. SPLITSTREE phylogenetic network of mitochondrial haplotypes. Tip labels are color coded as *P.p. profundus* (purple), *P.p. dubius* (blue), and *P. youngi* (green).

Color Analysis: Color morphs (i.e. species) appear to comprise distinct entities. Further, the “lateral” and “anterior” elytral areas best differentiate these morphs based on p-values from statistical tests (Figure 10). The lateral region separated *P.p. profundus* and *P. youngi* significantly, while the anterior region separated *P. youngi* and *P.p. dubius* as well as *P.p. profundus* and *P. youngi* significantly. PCA clustered taxa based on the four dorsal regions, and PERMANOVA showed differences in taxa through statistical analysis. Based on this test, *P. youngi* and *P.p. dubius*

are separated significantly based on color morphs, as well as *P.p. profundus* and *P. youngi* (Figure 11).

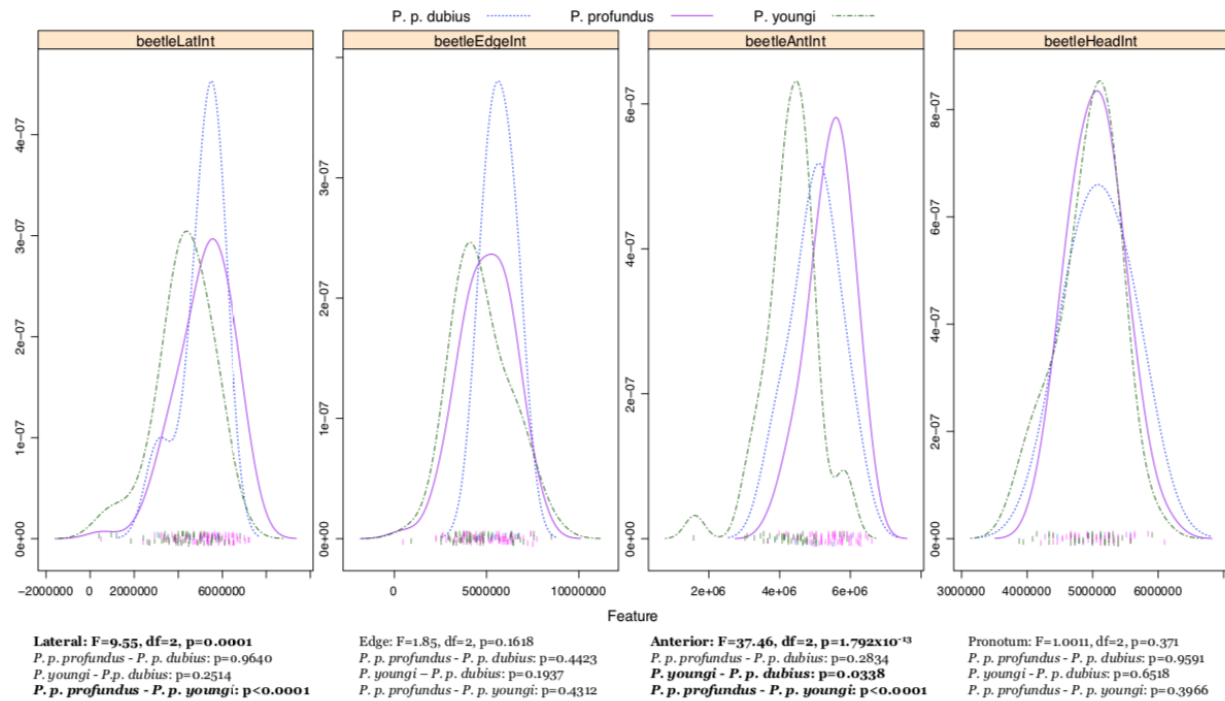


Figure 10. Color frequency distributions (as integers) for the four dorsal regions and associated statistical results. Taxa were compared via ANOVAs and results were investigated using Tukey's *post-hoc* test. Significant results are bolded.

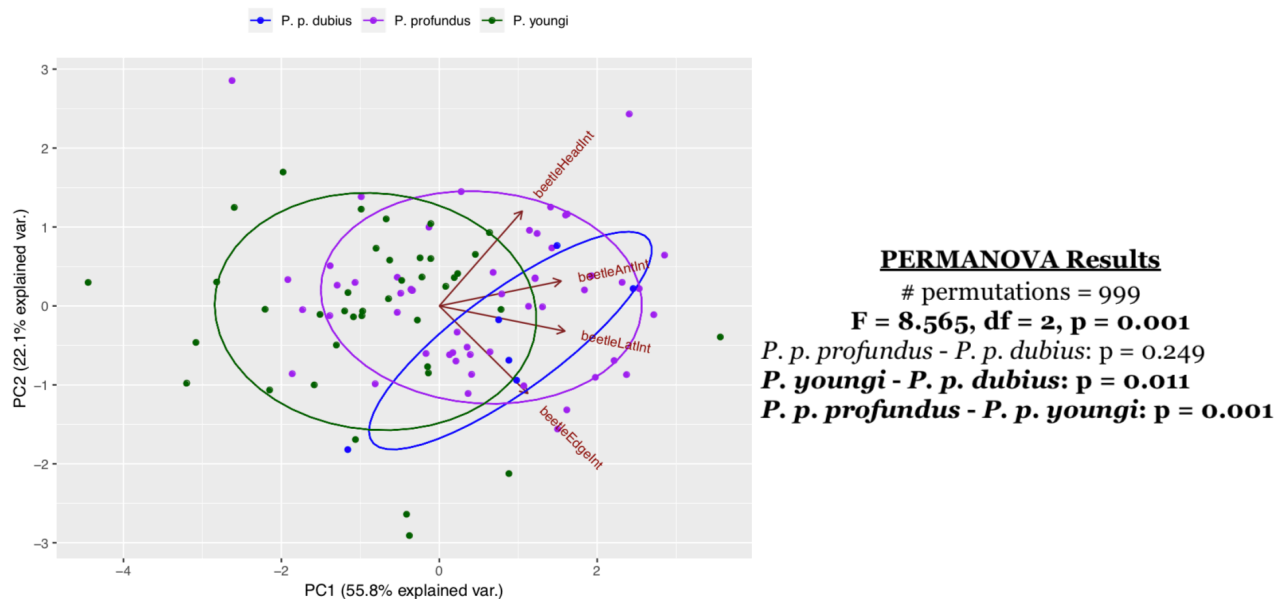


Figure 11. Principle component analysis and PERMANOVA results for color measurements. Significant results are bolded.

Discussion

Although the mitochondrial gene analysis did not support the monophyly of the three taxa of *Pelotrupes*, their color morphs appear to comprise distinct entities. While these taxa are not supported genetically, they are supported through their color morphs. Interestingly, these beetles are burrowers and spend their time above ground in darkness (Howden, 1955). Further, the iridescent coloration that distinguishes them is structural, meaning it is caused by light rather than pigment. With this knowledge, it is interesting to see how strongly color separates these burrowing beetles, while mitochondrial genetic analysis does not. Although these beetles inhabit geographically distinct regions, there is no phylogenetic structure defining the distinct species. These inconsistencies and the results from the color analysis raise a new question: why do discrete color morphs exist despite apparent presence of gene flow?

Conclusion

The analysis of genetic relationships among defined species and subspecies of *Peltotrupes* showed that these taxa are not monophyletic, as was previously thought. However, distinct coloration does appear to distinguish them, particularly from the “lateral” and “anterior” elytral regions. The ANOVA and the PERMANOVA results support significant differences between *P.p. profundus* and *P. youngi* as well as between *P.p. dubius* and *P. youngi*. However, no significant differences were observed between the subspecies *P.p. profundus* and *P.p. dubius*. Differences in the genetic and morphological results lead us to believe that an ecological role is at play in maintaining the distinct color morphs.

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Appendix

Peltotrupes Color Measurement Protocol

****DO NOT TAKE MEASUREMENTS WHERE THERE IS A GLARE ON THE SPECIMEN****

- Step 1. Open GIMP.
- Step 2. Select a file to edit.
- Select “okay” for TIFF image Message pop up and select “convert” for Convert to RGB Working Space.
- Step 3. Once file is open, zoom in on location being measured (*command, scroll*) and select the “Color Picker Tool”. **Note: this looks like an eyedropper.**
- After selecting tool, mark “sample average” box of the bottom left toolbar, so the radius can be adjusted.
 - Mode should be on “set foreground color”.
- Step 4. Adjust radius to 50 and select an area of grey near specimen, but be sure the area is consistent without any difference in coloration or brightness. **Note: the best place is usually right above the head, to the right of the first leg.**
- To get the RGB data of the area, use the method: “shift key, command key, and click” and the values will appear in a small box.
 - Record grey measurements for red, green, and blue in spreadsheet for lateral, edge, anterior, and head.
- Step 5. Adjust radius to 10 and select area of the elytrum along the edge where color is most reflective/colorful. Color is most consistent/standard in the thickest part of the lateral area.
- There will be two measurements for this area-one along the lateral (inside ridge) and one along the very edge of the elytrum. **Note: there is a definite “ridge” that separates the lateral and edge measurements.**
 - Collect the RGB values for both areas of measurement and record them in the spreadsheet (one under “lateral” and the other under “edge”).
 - **Note: the radius may need to be adjusted to a smaller size in order to get a clean measurement of the edge. **MAKE SURE TO GET A CLEAN, CONSISTENT SAMPLE****
- Step 6. Keep radius at 10 and select area at the top of the elytrum, right as it is joined to the thorax.
- Pick a spot where the color is reflective and colorful and record the RGB values in the spreadsheet

- **Note: count 4-6 lines left of the center of the elytra and get a measurement in this vicinity.**

Step 7. With the radius still at 10, select area at the base of the head (right where the head and thorax connect).

- Pick a spot (without glare) between the center of the head and the left eye and record RGB values into spreadsheet.

Step 8. To measure the data in Excel, type “=” and then the cell number of the color of the sample and divide that by the cell number of the same color for the grey measurement. Once doing the first box for the correction, you will be able to shift down the same formula.

- **EXAMPLE:** To get the corrected value for each RGB value, if the R value for grey measurement is 205 (and in cell B3), and the R value for the “lateral” measurement is 132 (and in cell C3), then in the cell for the corrected value, write “=C3/B3” and “enter”.
- **NOTE:** this should give a small decimal value→ once getting this value, drag the bottom right corner of the cell across to get the rest of the Green and Blue values, and then drag down to do the same for the remaining samples of each specimen.
- Do the same thing for the edge, anterior, and head recorded measurements.

	Grey - Lateral			Sample - Lateral			Corrected RGB - Lateral			Standardized RGB - Lateral			Species
Sample	R	G	B	R	G	B	R	G	B	R	G	B	
P120	211	203	192	84	63	91	0.39810427	0.31034483	0.47395833	90	67.5	97.5	<i>P. profundus</i>
P142	202	193	181	65	97	114	0.32178218	0.50259067	0.62983425	60.0543478	89.6195652	105.326087	<i>P. youngi</i>
P002	201	190	178	35	81	104	0.17412935	0.42631579	0.58426966	40.5681818	93.8863636	120.545455	<i>P. youngi</i>
P008	202	192	181	66	47	84	0.32673267	0.24479167	0.4640884	85.4314721	60.8375635	108.730964	<i>P. youngi</i>
P009	179	165	144	30	58	82	0.16759777	0.35151515	0.56944444	45	87	123	<i>P. youngi</i>
P010	194	183	169	56	86	129	0.28865979	0.46994536	0.76331361	52.6937269	80.9225092	121.383764	<i>P. youngi</i>
P011	188	177	161	32	79	86	0.17021277	0.44632768	0.53416149	41.4213198	102.258883	111.319797	<i>P. youngi</i>
P012	204	191	175	89	67	87	0.43627451	0.35078534	0.49714286	93.3950617	70.308642	91.2962963	<i>P. youngi</i>
P014	200	188	174	87	116	114	0.435	0.61702128	0.65517241	69.9842271	93.3123028	91.70347	<i>P. youngi</i>
P015	193	183	170	64	106	122	0.33160622	0.57923497	0.71764706	55.890411	92.5684932	106.541096	<i>P. youngi</i>
P016	204	192	178	59	70	92	0.28921569	0.36458333	0.51685393	68.0769231	80.7692308	106.153846	<i>P. youngi</i>
P017	208	197	183	79	63	96	0.37980769	0.31979695	0.52459016	84.6428571	67.5	102.857143	<i>P. youngi</i>
P018	208	197	183	78	100	121	0.375	0.50761421	0.66120219	66.5217391	85.2842809	103.19398	<i>P. youngi</i>
P019	206	196	183	70	110	108	0.33980583	0.56122449	0.59016393	61.9791667	97.3958333	95.625	<i>P. youngi</i>
P020	209	198	184	102	86	62	0.48803828	0.43434343	0.33695652	104.04	87.72	63.24	<i>P. youngi</i>
P021	199	186	169	53	62	85	0.26633166	0.33333333	0.50295858	67.575	79.05	108.375	<i>P. youngi</i>
P022	204	192	178	67	54	110	0.32843137	0.28125	0.61797753	73.961039	59.6103896	121.428571	<i>P. profundus</i>
P031	200	187	172	58	75	80	0.29	0.40106952	0.46511628	69.4366197	89.7887324	95.7746479	<i>P. youngi</i>
P032	204	193	179	73	120	122	0.35784314	0.62176166	0.68156425	59.0952381	97.1428571	98.7619048	<i>P. youngi</i>
P034	204	192	176	77	83	127	0.37745098	0.43229167	0.72159091	68.4146341	73.7456446	112.839721	<i>P. profundus</i>
P040	202	194	184	16	108	145	0.07920792	0.55670103	0.78804348	15.1672862	102.379182	137.453532	<i>P. youngi</i>
P049	209	200	188	77	67	102	0.36842105	0.335	0.54255319	79.8170732	69.4512195	105.731707	<i>P. youngi</i>
P050	203	193	181	30	74	98	0.14778325	0.38341969	0.54143646	37.8712871	93.4158416	123.712871	<i>P. youngi</i>
P051	203	193	181	19	115	131	0.09359606	0.59585492	0.72375691	18.2830189	110.660377	126.056604	<i>P. youngi</i>
P053	202	191	179	52	76	115	0.25742574	0.39790576	0.6424581	54.5679012	79.7530864	120.679012	<i>P. youngi</i>
P054	204	194	182	53	57	100	0.25980392	0.29381443	0.54945055	64.3571429	69.2142857	121.428571	<i>P. youngi</i>
P055	201	195	182	85	134	145	0.42288557	0.68717949	0.7967033	59.5467033	93.8736264	101.57967	<i>P. youngi</i>
P056	201	190	178	84	62	83	0.41791045	0.32631579	0.46629213	93.5371179	69.0393013	92.4235808	<i>P. profundus</i>
P057	202	193	182	72	46	76	0.35643564	0.23834197	0.41758242	94.6391753	60.4639175	99.8969072	<i>P. profundus</i>

P058	208	198	186	85	60	80	0.40865385	0.3030303	0.43010753	96.3333333	68	90.6666667	<i>P. profundus</i>
P059	210	201	190	93	59	85	0.44285714	0.29353234	0.44736842	100.063291	63.4810127	91.4556962	<i>P. profundus</i>
P060	212	202	192	86	56	85	0.40566038	0.27722772	0.44270833	96.6079295	62.907489	95.4845815	<i>P. profundus</i>
P061	208	202	191	60	81	123	0.28846154	0.4009901	0.64397906	57.9545455	78.2386364	118.806818	<i>P. profundus</i>
P062	217	208	198	61	109	158	0.28110599	0.52403846	0.7979798	47.4237805	84.7408537	122.835366	<i>P. p. dubius</i>
P064	219	210	201	109	67	96	0.49771689	0.31904762	0.47761194	102.1875	62.8125	90	<i>P. profundus</i>
P071	215	207	197	118	93	110	0.54883721	0.44927536	0.55837563	93.7383178	73.8785047	87.3831776	<i>P. profundus</i>
P074	211	203	192	68	61	103	0.32227488	0.30049261	0.53645833	74.7413793	67.0474138	113.211207	<i>P. profundus</i>
P075	207	202	192	87	86	117	0.42028986	0.42574257	0.609375	76.5	75.6206897	102.87931	<i>P. profundus</i>
P078	207	198	188	83	68	103	0.40096618	0.34343434	0.54787234	83.3267717	68.2677165	103.405512	<i>P. profundus</i>
P079	212	204	195	74	61	81	0.3490566	0.29901961	0.41538462	87.3611111	72.0138889	95.625	<i>P. profundus</i>
P080	210	201	192	103	65	102	0.49047619	0.32338308	0.53125	97.2777778	61.3888889	96.3333333	<i>P. profundus</i>
P081	202	193	178	85	69	87	0.42079208	0.35751295	0.48876404	89.9377593	73.0082988	92.0539419	<i>P. profundus</i>
P082	207	200	187	122	101	133	0.58937198	0.505	0.71122995	87.3876404	72.3455056	95.2668539	<i>P. profundus</i>
P083	210	200	189	83	59	114	0.3952381	0.295	0.6031746	82.6757813	58.7695313	113.554688	<i>P. profundus</i>
P086	212	202	191	108	71	90	0.50943396	0.35148515	0.47120419	102.379182	67.3048327	85.3159851	<i>P. profundus</i>
P087													<i>P. profundus</i>
P088	194	185	173	98	74	117	0.50515464	0.4	0.67630058	86.4705882	65.2941176	103.235294	<i>P. profundus</i>
P089	195	179	166	82	75	115	0.42051282	0.41899441	0.69277108	76.875	70.3125	107.8125	<i>P. profundus</i>
P091	193	184	171	93	60	99	0.48186528	0.32608696	0.57894737	94.1071429	60.7142857	100.178571	<i>P. profundus</i>
P093	194	181	167	86	62	73	0.44329897	0.34254144	0.43712575	99.2307692	71.5384615	84.2307692	<i>P. profundus</i>
P094	198	188	175	83	47	81	0.41919192	0.25	0.46285714	100.308057	56.8009479	97.8909953	<i>P. profundus</i>
P095	199	185	175	56	77	91	0.28140704	0.41621622	0.52	63.75	87.65625	103.59375	<i>P. profundus</i>
P096	190	180	167	91	65	66	0.47894737	0.36111111	0.39520958	104.527027	74.6621622	75.8108108	<i>P. profundus</i>
P097	190	176	149	59	81	114	0.31052632	0.46022727	0.76510067	59.2322835	81.3188976	114.448819	<i>P. profundus</i>
P098	179	168	150	43	46	91	0.24022346	0.27380952	0.60666667	60.9166667	65.1666667	128.916667	<i>P. profundus</i>
P099	201	191	178	69	57	111	0.34328358	0.29842932	0.62359551	74.2405063	61.3291139	119.43038	<i>P. profundus</i>
P100	175	165	154	44	66	111	0.25142857	0.4	0.72077922	50.7692308	76.1538462	128.076923	<i>P. youngi</i>
P101	196	187	176	61	103	101	0.31122449	0.55080214	0.57386364	58.6981132	99.1132075	97.1886792	<i>P. youngi</i>
P102	193	183	172	8	133	159	0.04145078	0.72677596	0.9244186	6.8	113.05	135.15	<i>P. youngi</i>
P103	185	172	157	69	96	96	0.37297297	0.55813953	0.61146497	67.4137931	93.7931034	93.7931034	<i>P. youngi</i>

P104	191	180	167	34	125	149	0.17801047	0.69444444	0.89221557	28.1493506	103.49026	123.36039	<i>P. youngi</i>
P105	191	180	168	38	76	84	0.19895288	0.42222222	0.5	48.9393939	97.8787879	108.181818	<i>P. youngi</i>
P114	185	173	161	39	64	82	0.21081081	0.3699422	0.50931677	53.7567568	88.2162162	113.027027	<i>P. profundus</i>
P116	185	174	162	64	45	68	0.34594595	0.25862069	0.41975309	92.2033898	64.8305085	97.9661017	<i>P. profundus</i>
P119	183	172	156	53	72	139	0.28961749	0.41860465	0.89102564	51.1931818	69.5454545	134.261364	<i>P. profundus</i>
P121	194	183	171	72	55	78	0.37113402	0.30054645	0.45614035	89.5609756	68.4146341	97.0243902	<i>P. profundus</i>
P122	185	175	163	99	79	118	0.53513514	0.45142857	0.72392638	85.2871622	68.0574324	101.655405	<i>P. profundus</i>
P123													<i>P. profundus</i>
P125	179	171	157	64	51	78	0.3575419	0.29824561	0.49681529	84.5595855	67.3834197	103.056995	<i>P. profundus</i>
P127	178	167	155	49	87	111	0.2752809	0.52095808	0.71612903	50.5870445	89.8178138	114.595142	<i>P. profundus</i>
P128	180	172	158	33	61	89	0.18333333	0.35465116	0.56329114	45.9836066	85	124.016393	<i>P. profundus</i>
P129													<i>P. profundus</i>
P130	199	187	175	108	96	134	0.54271357	0.51336898	0.76571429	81.4792899	72.4260355	101.094675	<i>P. profundus</i>
P134	182	171	158	95	73	122	0.52197802	0.42690058	0.7721519	83.5344828	64.1896552	107.275862	<i>P. profundus</i>
P135	194	184	173	46	116	147	0.2371134	0.63043478	0.84971098	37.961165	95.7281553	121.31068	<i>P. profundus</i>
P136	195	186	174	102	161	183	0.52307692	0.8655914	1.05172414	58.3183857	92.0515695	104.630045	<i>P. profundus</i>
P137	201	192	179	9	101	160	0.04477612	0.52604167	0.89385475	8.5	95.3888889	151.111111	<i>P. profundus</i>
P138	199	188	176	65	102	155	0.32663317	0.54255319	0.88068182	51.4751553	80.7763975	122.748447	<i>P. profundus</i>
P139	210	200	186	77	64	97	0.36666667	0.32	0.52150538	82.5	68.5714286	103.928571	<i>P. profundus</i>
P140	200	189	177	59	131	171	0.295	0.69312169	0.96610169	41.6759003	92.534626	120.789474	<i>P. profundus</i>
P141	201	189	174	68	99	122	0.33830846	0.52380952	0.70114943	60	87.3529412	107.647059	<i>P. profundus</i>
P143	214	207	198	75	69	123	0.35046729	0.33333333	0.62121212	71.6292135	65.8988764	117.47191	<i>P. profundus</i>
P144	199	188	175	55	52	104	0.27638191	0.27659574	0.59428571	66.4691943	62.8436019	125.687204	<i>P. profundus</i>
P145	203	196	184	77	65	113	0.37931034	0.33163265	0.61413043	77	65	113	<i>P. profundus</i>
P147	205	194	182	72	111	116	0.35121951	0.57216495	0.63736264	61.4046823	94.6655518	98.9297659	<i>P. youngi</i>
P149	209	203	194	74	59	107	0.35406699	0.29064039	0.55154639	78.625	62.6875	113.6875	<i>P. profundus</i>
P152	207	201	191	86	121	155	0.41545894	0.60199005	0.81151832	60.5801105	85.2348066	109.185083	<i>P. profundus</i>
P154	180	169	156	39	31	87	0.21666667	0.18343195	0.55769231	63.343949	50.3503185	141.305732	<i>P. profundus</i>
P155	185	174	161	58	44	72	0.31351351	0.25287356	0.44720497	85	64.4827586	105.517241	<i>P. profundus</i>
P026	173	166	154	45	41	73	0.26011561	0.24698795	0.47402597	72.1698113	65.754717	117.075472	<i>P. profundus</i>
P025	205	199	187	129	122	169	0.62926829	0.61306533	0.90374332	78.3214286	74.0714286	102.607143	<i>P. profundus</i>

P159	181	173	163	70	72	114	0.38674033	0.41618497	0.6993865	69.7265625	71.71875	113.554688	<i>P. profundus</i>
P107	185	174	161	71	59	74	0.38378378	0.33908046	0.45962733	88.75	73.75	92.5	<i>P. p. dubius</i>
P157	172	163	148	51	60	110	0.29651163	0.36809816	0.74324324	58.8461538	69.2307692	126.923077	<i>P. profundus</i>
P005	175	168	155	39	54	101	0.22285714	0.32142857	0.6516129	51.2628866	70.9793814	132.757732	<i>P. profundus</i>
P004	180	170	156	57	39	67	0.31666667	0.22941176	0.42948718	89.1717791	61.0122699	104.815951	<i>P. profundus</i>
P166	181	171	156	55	36	66	0.3038674	0.21052632	0.42307692	89.3312102	58.4713376	107.197452	
P167	183	172	159	59	50	77	0.32240437	0.29069767	0.48427673	80.8870968	68.5483871	105.564516	
P027	177	169	156	73	72	130	0.41242938	0.4260355	0.83333333	67.6909091	66.7636364	120.545455	<i>P. profundus</i>
P148	210	204	192	48	35	85	0.22857143	0.17156863	0.44270833	72.8571429	53.125	129.017857	<i>P. profundus</i>
P158	203	195	183	100	112	119	0.49261084	0.57435897	0.65027322	77.0392749	86.2839879	91.6767372	<i>P. youngi</i>

	Grey- Edge Elytrum			Sample - Edge Elytrum			Corrected RGB - Edge Elytrum			Standardized - RGB		
Sample	R	G	B	R	G	B	R	G	B	R	G	B
P120	211	203	192	79	66	91	0.37440758	0.32512315	0.47395833	85.3601695	71.3135593	98.3262712
P142	202	193	181	133	175	142	0.65841584	0.90673575	0.78453039	75.3666667	99.1666667	80.4666667
P002	201	190	178	67	158	154	0.33333333	0.83157895	0.86516854	45.0791557	106.306069	103.614776
P008	202	192	181	64	86	118	0.31683168	0.44791667	0.6519337	60.8955224	81.8283582	112.276119
P009	179	165	144	69	155	152	0.38547486	0.93939394	1.05555556	46.7952128	105.119681	103.085106
P010	194	183	169	103	71	89	0.53092784	0.38797814	0.52662722	99.8669202	68.8403042	86.2927757
P011	188	177	161	68	94	88	0.36170213	0.53107345	0.54658385	69.36	95.88	89.76
P012	204	191	175	121	78	98	0.59313725	0.40837696	0.56	103.888889	66.969697	84.1414141
P014	200	188	174	133	86	71	0.665	0.45744681	0.40804598	116.948276	75.6206897	62.4310345
P015	193	183	170	119	172	165	0.61658031	0.93989071	0.97058824	66.5460526	96.1842105	92.2697368
P016	204	192	178	133	76	81	0.65196078	0.39583333	0.45505618	116.948276	66.8275862	71.2241379
P017	208	197	183	111	151	179	0.53365385	0.76649746	0.97814208	64.1836735	87.3129252	103.503401
P018	208	197	183	98	161	185	0.47115385	0.81725888	1.01092896	56.2837838	92.4662162	106.25
P019	206	196	183	147	191	164	0.71359223	0.9744898	0.89617486	74.6713147	97.0219124	83.3067729
P020	209	198	184	142	74	61	0.67942584	0.37373737	0.33152174	130.722022	68.1227437	56.1552347
P021	199	186	169	55	88	95	0.27638191	0.47311828	0.56213018	58.9285714	94.2857143	101.785714
P022	204	192	178	70	146	216	0.34313725	0.76041667	1.21348315	41.3194444	86.1805556	127.5

P031	200	187	172	86	116	104	0.43	0.62032086	0.60465116	71.6666667	96.6666667	86.6666667
P032	204	193	179	129	160	152	0.63235294	0.82901554	0.84916201	74.5918367	92.5170068	87.8911565
P034	204	192	176	127	106	125	0.62254902	0.55208333	0.71022727	90.4608939	75.5027933	89.0363128
P040	202	194	184	70	192	193	0.34653465	0.98969072	1.04891304	39.2307692	107.604396	108.164835
P049	209	200	188	43	110	126	0.20574163	0.55	0.67021277	39.3010753	100.537634	115.16129
P050	203	193	181	107	174	168	0.5270936	0.9015544	0.9281768	60.7683742	98.8195991	95.4120267
P051	203	193	181	96	191	178	0.4729064	0.98963731	0.98342541	52.6451613	104.741935	97.6129032
P053	202	191	179	70	149	145	0.34653465	0.78010471	0.81005587	49.0384615	104.381868	101.57967
P054	204	194	182	99	60	80	0.48529412	0.30927835	0.43956044	105.627615	64.0167364	85.3556485
P055	201	195	182	138	191	164	0.68656716	0.97948718	0.9010989	71.3793103	98.7931034	84.8275862
P056	201	190	178	125	62	89	0.62189055	0.32631579	0.5	115.48913	57.2826087	82.2282609
P057	202	193	182	78	101	132	0.38613861	0.52331606	0.72527473	63.9549839	82.8135048	108.231511
P058	208	198	186	112	68	95	0.53846154	0.34343434	0.51075269	103.854545	63.0545455	88.0909091
P059	210	201	190	120	128	170	0.57142857	0.63681592	0.89473684	73.2057416	78.0861244	103.708134
P060	212	202	192	90	67	80	0.4245283	0.33168317	0.41666667	96.835443	72.0886076	86.0759494
P061	208	202	191	136	108	124	0.65384615	0.53465347	0.64921466	94.2391304	74.8369565	85.923913
P062	217	208	198	147	99	121	0.67741935	0.47596154	0.61111111	102.138965	68.7874659	84.0735695
P064	219	210	201	108	129	193	0.49315068	0.61428571	0.960199	64.0465116	76.5	114.453488
P071	215	207	197	184	115	123	0.85581395	0.55555556	0.62436548	111.184834	69.4905213	74.3246445
P074	211	203	192	108	81	122	0.51184834	0.39901478	0.63541667	88.5530547	66.414791	100.032154
P075	207	202	192	125	71	73	0.60386473	0.35148515	0.38020833	118.494424	67.3048327	69.2007435
P078	207	198	188	92	123	185	0.44444444	0.62121212	0.98404255	58.65	78.4125	117.9375
P079	212	204	195	127	177	191	0.5990566	0.86764706	0.97948718	65.4242424	91.1818182	98.3939394
P080	210	201	192	88	143	201	0.41904762	0.71144279	1.046875	51.9444444	84.4097222	118.645833
P081	202	193	178	116	80	107	0.57425743	0.41450777	0.6011236	97.6237624	67.3267327	90.049505
P082	207	200	187	104	142	211	0.50241546	0.71	1.12834225	58.0306346	79.2341357	117.73523
P083	210	200	189	133	145	185	0.63333333	0.725	0.97883598	73.25054	79.8596112	101.889849
P086	212	202	191	177	134	176	0.83490566	0.66336634	0.92146597	92.6796715	70.164271	92.1560575
P087												
P088	194	185	173	84	99	181	0.43298969	0.53513514	1.04624277	58.8461538	69.3543956	126.799451
P089	195	179	166	101	93	162	0.51794872	0.51955307	0.97590361	72.3455056	66.6151685	116.039326

P091	193	184	171	126	84	135	0.65284974	0.45652174	0.78947368	93.1304348	62.0869565	99.7826087
P093	194	181	167	106	69	125	0.54639175	0.38121547	0.74850299	90.1	58.65	106.25
P094	198	188	175	53	117	140	0.26767677	0.62234043	0.8	43.5967742	96.2419355	115.16129
P095	199	185	175	66	133	171	0.33165829	0.71891892	0.97714286	45.4864865	91.6621622	117.851351
P096	190	180	167	106	78	108	0.55789474	0.43333333	0.64670659	92.5684932	68.1164384	94.3150685
P097	190	176	149	59	81	114	0.31052632	0.46022727	0.76510067	59.2322835	81.3188976	114.448819
P098	179	168	150	57	84	133	0.31843575	0.5	0.88666667	53.0474453	78.1751825	123.777372
P099	201	191	178	100	118	180	0.49751244	0.61780105	1.01123596	64.0703518	75.6030151	115.326633
P100	175	165	154	18	136	165	0.10285714	0.82424242	1.07142857	14.3887147	108.714734	131.896552
P101	196	187	176	81	158	139	0.41326531	0.84491979	0.78977273	54.6428571	106.587302	93.7698413
P102	193	183	172	81	165	152	0.41968912	0.90163934	0.88372093	51.8969849	105.71608	97.3869347
P103	185	172	157	64	103	91	0.34594595	0.59883721	0.57961783	63.255814	101.802326	89.9418605
P104	191	180	167	53	100	104	0.27748691	0.55555556	0.62275449	52.5875486	99.2217899	103.190661
P105	191	180	168	57	112	124	0.29842932	0.62222222	0.73809524	49.6075085	97.4744027	107.918089
P114	185	173	161	68	84	85	0.36756757	0.48554913	0.52795031	73.164557	90.3797468	91.4556962
P116	185	174	162	80	42	57	0.43243243	0.24137931	0.35185185	113.96648	59.8324022	81.2011173
P119	183	172	156	111	78	97	0.60655738	0.45348837	0.62179487	98.9685315	69.5454545	86.486014
P121	194	183	171	111	152	153	0.57216495	0.83060109	0.89473684	68.0408654	93.1730769	93.7860577
P122	185	175	163	104	81	117	0.56216216	0.46285714	0.71779141	87.8145695	68.3940397	98.7913907
P123												
P124	179	170	157	74	103	143	0.41340782	0.60588235	0.91082803	58.96875	82.078125	113.953125
P125	179	171	157	72	104	144	0.40223464	0.60818713	0.91719745	57.375	82.875	114.75
P127	178	167	155	109	156	146	0.61235955	0.93413174	0.94193548	67.6277372	96.7883212	90.5839416
P128	180	172	158	90	122	120	0.5	0.70930233	0.75949367	69.126506	93.7048193	92.1686747
P129	197	187	175	74	126	144	0.37563452	0.67379679	0.82285714	54.8546512	93.4011628	106.744186
P130	199	187	175	123	140	178	0.61809045	0.7486631	1.01714286	71.122449	80.952381	102.92517
P134	182	171	158	118	84	128	0.64835165	0.49122807	0.81012658	91.1818182	64.9090909	98.9090909
P135	194	184	173	114	167	182	0.58762887	0.9076087	1.05202312	62.7861771	91.9762419	100.237581
P136	195	186	174	112	181	195	0.57435897	0.97311828	1.12068966	58.5245902	94.579918	101.895492
P137	201	192	179	9	142	167	0.04477612	0.73958333	0.93296089	7.21698113	113.867925	133.915094
P138	199	188	176	44	117	158	0.22110553	0.62234043	0.89772727	35.1724138	93.5266458	126.30094

P139	210	200	186	88	143	173	0.41904762	0.715	0.93010753	55.5445545	90.259901	109.195545
P140	200	189	177	81	163	165	0.405	0.86243386	0.93220339	50.5012225	101.625917	102.872861
P141	201	189	174	101	87	101	0.50248756	0.46031746	0.58045977	89.1176471	76.7647059	89.1176471
P143	214	207	198	111	133	173	0.51869159	0.64251208	0.87373737	67.8776978	81.3309353	105.791367
P144	199	188	175	47	106	141	0.2361809	0.56382979	0.80571429	40.7653061	91.9387755	122.295918
P145	203	196	184	103	99	149	0.50738916	0.50510204	0.80978261	74.8290598	71.9230769	108.247863
P147	205	194	182	99	154	149	0.48292683	0.79381443	0.81868132	62.7985075	97.6865672	94.5149254
P149	209	203	194	99	108	191	0.47368421	0.5320197	0.98453608	63.4296482	69.1959799	122.374372
P152	207	201	191	114	142	170	0.55072464	0.70646766	0.89005236	68.2394366	85	101.760563
P154	180	169	156	30	68	117	0.16666667	0.40236686	0.75	35.5813953	80.6511628	138.767442
P155	185	174	161	97	55	78	0.52432432	0.31609195	0.48447205	107.543478	60.9782609	86.4782609
P026	173	166	154	75	100	147	0.43352601	0.60240964	0.95454545	59.3944099	79.1925466	116.413043
P025	205	199	187	127	150	189	0.6195122	0.75376884	1.01069519	69.4957082	82.0815451	103.422747
P159	181	173	163	112	82	109	0.61878453	0.47398844	0.66871166	94.2574257	69.009901	91.7326733
P107	185	174	161	105	84	96	0.56756757	0.48275862	0.59627329	93.9473684	75.1578947	85.8947368
P157	172	163	148	91	127	151	0.52906977	0.7791411	1.02027027	62.8861789	87.7642276	104.349593
P005	175	168	155	53	91	123	0.30285714	0.54166667	0.79354839	50.6179775	86.9101124	117.47191
P004	180	170	156	79	98	158	0.43888889	0.57647059	1.01282051	60.1343284	74.5970149	120.268657
P166	181	171	156	46	56	107	0.25414365	0.32748538	0.68589744	56.1244019	68.3253589	130.550239
P167	183	172	159	99	75	102	0.54098361	0.43604651	0.64150943	91.4673913	69.2934783	94.2391304
P027	177	169	156	117	95	127	0.66101695	0.56213018	0.81410256	88.0088496	71.460177	95.5309735
P148	210	204	192	106	94	177	0.5047619	0.46078431	0.921875	71.6976127	63.5809019	119.721485
P158	203	195	183	121	143	153	0.59605911	0.73333333	0.83606557	73.9928058	87.4460432	93.5611511

	Grey - Anterior			Sample - Anterior			Corrected RGB - Anterior			Standardized RGB - Anterior		
Sample	R	G	B	R	G	B	R	G	B	R	G	B
P120	211	203	192	132	111	143	0.62559242	0.54679803	0.74479167	87.2020725	73.3290155	94.4689119
P142	202	193	181	142	168	160	0.7029703	0.87046632	0.8839779	77.0425532	91.1489362	86.8085106
P002	201	190	178	60	124	140	0.29850746	0.65263158	0.78651685	47.2222222	97.5925926	110.185185
P008	202	192	181	86	110	137	0.42574257	0.57291667	0.75690608	65.8558559	84.2342342	104.90991
P009	179	165	144	51	119	110	0.2849162	0.72121212	0.76388889	46.4464286	108.375	100.178571

P010	194	183	169	97	126	135	0.5	0.68852459	0.79881657	69.0921788	89.7486034	96.1592179
P011	188	177	161	91	123	135	0.48404255	0.69491525	0.83850932	66.4899713	89.8710602	98.6389685
P012	204	191	175	101	104	132	0.49509804	0.54450262	0.75428571	76.4243323	78.694362	99.8813056
P014	200	188	174	74	108	115	0.37	0.57446809	0.66091954	63.5353535	92.7272727	98.7373737
P015	193	183	170	103	156	160	0.53367876	0.85245902	0.94117647	62.6849642	94.9403341	97.3747017
P016	204	192	178	90	146	147	0.44117647	0.76041667	0.8258427	59.921671	97.2062663	97.8720627
P017	208	197	183	119	143	176	0.57211538	0.72588832	0.96174863	69.2808219	83.2534247	102.465753
P018	208	197	183	85	128	161	0.40865385	0.64974619	0.87978142	57.9545455	87.2727273	109.772727
P019	206	196	183	157	205	193	0.76213592	1.04591837	1.05464481	72.1351351	94.1891892	88.6756757
P020	209	198	184	100	93	98	0.4784689	0.46969697	0.5326087	87.628866	81.4948454	85.8762887
P021	199	186	169	87	134	133	0.43718593	0.72043011	0.78698225	62.6694915	96.5254237	95.8050847
P022	204	192	178	95	89	122	0.46568627	0.46354167	0.68539326	79.1666667	74.1666667	101.666667
P031	200	187	172	87	110	114	0.435	0.58823529	0.6627907	71.3344051	90.192926	93.4726688
P032	204	193	179	76	122	120	0.37254902	0.63212435	0.67039106	60.9433962	97.8301887	96.2264151
P034	204	192	176	126	105	122	0.61764706	0.546875	0.69318182	91.01983	75.8498584	88.1303116
P040	202	194	184	107	157	164	0.52970297	0.80927835	0.89130435	63.75	93.5397196	97.7102804
P049	209	200	188	130	165	157	0.62200957	0.825	0.83510638	73.340708	93.0862832	88.5730088
P050	203	193	181	58	99	117	0.28571429	0.51295337	0.64640884	53.9781022	92.1350365	108.886861
P051	203	193	181	103	181	185	0.50738916	0.93782383	1.02209945	56.0021322	98.4115139	100.586354
P053	202	191	179	117	176	183	0.57920792	0.92146597	1.02234637	62.6785714	94.2857143	98.0357143
P054	204	194	182	74	126	149	0.3627451	0.64948454	0.81868132	54.0687679	92.0630372	108.868195
P055	201	195	182	100	126	141	0.49751244	0.64615385	0.77472527	69.4822888	87.5476839	97.9700272
P056	201	190	178	95	101	132	0.47263682	0.53157895	0.74157303	73.8567073	78.5213415	102.621951
P057	202	193	182	100	89	109	0.4950495	0.4611399	0.5989011	85.5704698	76.1577181	93.2718121
P058	208	198	186	112	93	119	0.53846154	0.46969697	0.63978495	88.1481481	73.1944444	93.6574074
P059	210	201	190	121	97	128	0.57619048	0.48258706	0.67368421	89.1763006	71.4884393	94.3352601
P060	212	202	192	147	120	137	0.69339623	0.59405941	0.71354167	92.7846535	75.7425743	86.4727723
P061	208	202	191	96	99	112	0.46153846	0.49009901	0.58638743	79.7394137	82.2312704	93.029316
P062	217	208	198	110	176	183	0.50691244	0.84615385	0.92424242	59.8081023	95.6929638	99.4989339
P064	219	210	201	129	117	178	0.5890411	0.55714286	0.88557214	77.5825472	70.365566	107.051887
P071	215	207	197	120	102	118	0.55813953	0.49275362	0.59898477	90	76.5	88.5

P074	211	203	192	152	155	201	0.72037915	0.7635468	1.046875	76.2992126	77.8051181	100.895669
P075	207	202	192	114	107	120	0.55072464	0.52970297	0.625	85.2492669	80.0146628	89.7360704
P078	207	198	188	129	106	119	0.62318841	0.53535354	0.63297872	92.9237288	76.3559322	85.720339
P079	212	204	195	128	107	132	0.60377358	0.5245098	0.67692308	88.9373297	74.346049	91.7166213
P080	210	201	192	151	97	157	0.71904762	0.48258706	0.81770833	95.0740741	61.0740741	98.8518519
P081	202	193	178	138	117	142	0.68316832	0.60621762	0.79775281	88.6397985	75.1511335	91.209068
P082	207	200	187	144	122	149	0.69565217	0.61	0.79679144	88.4819277	74.9638554	91.5542169
P083	210	200	189	77	55	98	0.36666667	0.275	0.51851852	85.3695652	60.9782609	108.652174
P086	212	202	191	146	117	159	0.68867925	0.57920792	0.83246073	88.2227488	70.6990521	96.0781991
P087												
P088	194	185	173	132	96	121	0.68041237	0.51891892	0.69942197	96.4469914	70.1432665	88.4097421
P089	195	179	166	88	74	111	0.45128205	0.41340782	0.6686747	82.1978022	69.1208791	103.681319
P091	193	184	171	115	85	105	0.59585492	0.46195652	0.61403509	96.147541	71.0655738	87.7868852
P093	194	181	167	139	107	125	0.71649485	0.59116022	0.74850299	95.5390836	73.5444744	85.916442
P094	198	188	175	115	94	146	0.58080808	0.5	0.83428571	82.6056338	67.5211268	104.873239
P095	199	185	175	112	100	132	0.56281407	0.54054054	0.75428571	83.0232558	74.127907	97.8488372
P096	190	180	167	135	104	117	0.71052632	0.57777778	0.7005988	96.6994382	74.494382	83.8061798
P097	190	176	149	112	97	150	0.58947368	0.55113636	1.00671141	79.5543175	68.8997214	106.545961
P098	179	168	150	163	140	182	0.91061453	0.83333333	1.21333333	85.7010309	73.6082474	95.6907216
P100	175	165	154	107	142	165	0.61142857	0.86060606	1.07142857	65.9057971	87.4637681	101.630435
P101	196	187	176	133	191	175	0.67857143	1.02139037	0.99431818	67.9659319	97.6052104	89.4288577
P102	193	183	172	26	118	125	0.13471503	0.64480874	0.72674419	24.6468401	111.858736	118.494424
P103	185	172	157	66	117	126	0.35675676	0.68023256	0.80254777	54.4660194	96.5533981	103.980583
P104	191	180	167	78	148	168	0.40837696	0.82222222	1.00598802	50.4822335	95.786802	108.730964
P105	191	180	168	90	132	104	0.47120419	0.73333333	0.61904762	70.398773	103.251534	81.3496933
P114	185	173	161	84	92	119	0.45405405	0.53179191	0.73913043	72.6101695	79.5254237	102.864407
P116	185	174	162	108	91	114	0.58378378	0.52298851	0.7037037	87.9872204	74.1373802	92.8753994
P119	183	172	156	106	132	149	0.57923497	0.76744186	0.95512821	69.8449612	86.9767442	98.1782946
P121	194	183	171	126	131	160	0.64948454	0.71584699	0.93567251	77.0503597	80.1079137	97.8417266
P122	185	175	163	118	108	130	0.63783784	0.61714286	0.79754601	84.5224719	77.3595506	93.1179775
P123												

P124												
P125												
P127	178	167	155	86	141	161	0.48314607	0.84431138	1.03870968	56.5206186	92.6675258	105.811856
P128	180	172	158	107	144	177	0.59444444	0.8372093	1.12025316	63.75	85.7943925	105.455607
P129												
P130	199	187	175	134	117	136	0.67336683	0.62566845	0.77714286	88.2945736	77.0930233	89.6124031
P134	182	171	158	124	111	134	0.68131868	0.64912281	0.84810127	85.6910569	76.7073171	92.601626
P135	194	184	173	120	162	191	0.6185567	0.88043478	1.10404624	64.6934461	87.3361522	102.970402
P136	195	186	174	112	120	145	0.57435897	0.64516129	0.83333333	75.7559682	81.1671088	98.0769231
P137	201	192	179	134	124	192	0.66666667	0.64583333	1.0726257	75.9333333	70.2666667	108.8
P138	199	188	176	132	127	172	0.66331658	0.67553191	0.97727273	78.0974478	75.1392111	101.763341
P139	210	200	186	133	104	145	0.63333333	0.52	0.77956989	88.7827225	69.4240838	96.7931937
P140	200	189	177	114	157	203	0.57	0.83068783	1.14689266	61.3291139	84.4620253	109.208861
P141	201	189	174	128	134	176	0.63681592	0.70899471	1.01149425	74.5205479	78.0136986	102.465753
P143	214	207	198	145	143	180	0.67757009	0.69082126	0.90909091	79.0064103	77.9166667	98.0769231
P144	199	188	175	132	158	195	0.66331658	0.84042553	1.11428571	69.4020619	83.0721649	102.525773
P145	203	196	184	114	104	164	0.56157635	0.53061224	0.89130435	76.0994764	69.4240838	109.47644
P147	205	194	182	132	190	204	0.64390244	0.97938144	1.12087912	63.9923954	92.1102662	98.8973384
P149	209	203	194	161	133	187	0.77033493	0.65517241	0.96391753	85.3534304	70.5093555	99.1372141
P152	207	201	191	127	146	198	0.61352657	0.72636816	1.03664921	68.7579618	79.044586	107.197452
P154	180	169	156	80	88	144	0.44444444	0.52071006	0.92307692	65.3846154	71.9230769	117.692308
P155	185	174	161	105	70	88	0.56756757	0.40229885	0.54658385	101.806084	67.8707224	85.3231939
P026	173	166	154	102	98	148	0.58959538	0.59036145	0.96103896	74.7413793	71.8103448	108.448276
P025	205	199	187	133	127	166	0.64878049	0.63819095	0.88770053	79.6126761	76.0211268	99.3661972
P159	181	173	163	123	159	185	0.67955801	0.91907514	1.13496933	67.1627409	86.8201285	101.017131
P107	185	174	161	99	110	127	0.53513514	0.63218391	0.78881988	75.1339286	83.4821429	96.3839286
P157	172	163	148	151	145	189	0.87790698	0.88957055	1.27702703	79.3917526	76.2371134	99.371134
P005	175	168	155	104	109	165	0.59428571	0.64880952	1.06451613	70.1587302	73.531746	111.309524
P004	180	170	156	114	104	128	0.63333333	0.61176471	0.82051282	84.017341	76.6473988	94.3352601
P166	181	171	156	81	82	129	0.44751381	0.47953216	0.82692308	70.7363014	71.609589	112.65411
P167	183	172	159	127	109	143	0.69398907	0.63372093	0.89937107	85.4485488	73.3377309	96.2137203

P027	177	169	156	111	89	129	0.62711864	0.52662722	0.82692308	86.0334347	68.9817629	99.9848024
P148	210	204	192	128	110	142	0.60952381	0.53921569	0.73958333	85.8947368	73.8157895	95.2894737
P158	203	195	183	141	177	179	0.69458128	0.90769231	0.97814208	72.3440644	90.8148893	91.8410463

	Grey - Head			Sample - Head			Corrected RGB - Head			Standardized RGB - Head		
Sample	R	G	B	R	G	B	R	G	B	R	G	B
P120	211	203	192	133	137	140	0.63033175	0.67487685	0.72916667	82.7195122	85.2073171	87.0731707
P142	202	193	181	129	151	128	0.63861386	0.78238342	0.70718232	80.625	94.375	80
P002	201	190	178	120	176	166	0.59701493	0.92631579	0.93258427	66.2337662	97.1428571	91.6233766
P008	202	192	181	85	130	141	0.42079208	0.67708333	0.77900552	60.8848315	93.1179775	100.997191
P009	179	165	144	74	122	104	0.41340782	0.73939394	0.72222222	62.9	103.7	88.4
P010	194	183	169	155	183	179	0.79896907	1	1.0591716	76.450677	90.2611219	88.2882012
P011	188	177	161	126	152	140	0.67021277	0.85875706	0.86956522	76.8660287	92.7272727	85.4066986
P012	204	191	175	94	123	137	0.46078431	0.64397906	0.78285714	67.7118644	88.6016949	98.6864407
P014	200	188	174	100	145	159	0.5	0.7712766	0.9137931	63.1188119	91.5222772	100.358911
P015	193	183	170	91	126	133	0.47150259	0.68852459	0.78235294	66.3	91.8	96.9
P016	204	192	178	82	102	129	0.40196078	0.53125	0.7247191	66.8051118	83.0990415	105.095847
P017	208	197	183	118	111	123	0.56730769	0.56345178	0.67213115	85.4829545	80.4119318	89.1051136
P018	208	197	183	164	194	157	0.78846154	0.98477157	0.8579235	81.2038835	96.0582524	77.7378641
P019	206	196	183	143	175	164	0.69417476	0.89285714	0.89617486	75.653527	92.5829876	86.7634855
P020	209	198	184	128	131	100	0.61244019	0.66161616	0.54347826	90.9192201	93.0501393	71.0306407
P021	199	186	169	90	109	112	0.45226131	0.58602151	0.66272189	73.7942122	89.3729904	91.8327974
P022	204	192	178	98	97	126	0.48039216	0.50520833	0.70786517	77.8504673	77.0560748	100.093458
P031	200	187	172	90	101	96	0.45	0.54010695	0.55813953	79.9651568	89.738676	85.2961672
P032	204	193	179	127	149	129	0.62254902	0.77202073	0.72067039	79.962963	93.8148148	81.2222222
P034	204	192	176	110	109	134	0.53921569	0.56770833	0.76136364	79.4617564	78.7393768	96.7988669
P040	202	194	184	124	153	134	0.61386139	0.78865979	0.72826087	76.9343066	94.9270073	83.1386861
P049												
P050	203	193	181	110	134	137	0.54187192	0.69430052	0.75690608	73.6220472	89.6850394	91.6929134
P051	203	193	181	115	140	147	0.56650246	0.7253886	0.8121547	72.9477612	88.8059701	93.2462687
P053	202	191	179	152	170	149	0.75247525	0.89005236	0.83240223	82.2929936	92.0382166	80.6687898

P054	204	194	182	121	132	129	0.59313725	0.68041237	0.70879121	80.7722513	88.1151832	86.1125654
P055	201	195	182	118	132	131	0.58706468	0.67692308	0.71978022	78.976378	88.3464567	87.6771654
P056	201	190	178	90	99	110	0.44776119	0.52105263	0.61797753	76.7558528	84.4314381	93.812709
P057	202	193	182	91	96	109	0.45049505	0.49740933	0.5989011	78.3952703	82.7027027	93.902027
P058	208	198	186	118	119	141	0.56730769	0.6010101	0.75806452	79.6031746	80.2777778	95.1190476
P059	210	201	190	125	114	153	0.5952381	0.56716418	0.80526316	81.3137755	74.1581633	99.5280612
P060	212	202	192	169	149	164	0.79716981	0.73762376	0.85416667	89.4087137	78.8278008	86.7634855
P061	208	202	191	97	114	113	0.46634615	0.56435644	0.59162304	76.3425926	89.7222222	88.9351852
P062	217	208	198	112	136	166	0.51612903	0.65384615	0.83838384	68.9855072	83.7681159	102.246377
P064	219	210	201	149	138	187	0.6803653	0.65714286	0.93034826	80.1582278	74.2405063	100.601266
P071	215	207	197	93	102	105	0.43255814	0.49275362	0.53299492	79.05	86.7	89.25
P074	211	203	192	123	143	158	0.58293839	0.7044335	0.82291667	73.9740566	86.0023585	95.0235849
P075	207	202	192	110	124	135	0.53140097	0.61386139	0.703125	76.0162602	85.6910569	93.2926829
P078	207	198	188	130	124	158	0.62801932	0.62626263	0.84042553	80.461165	76.7475728	97.7912621
P079	212	204	195	142	127	155	0.66981132	0.62254902	0.79487179	85.4009434	76.379717	93.2193396
P080	210	201	192	162	128	153	0.77142857	0.63681592	0.796875	93.2505643	73.6794582	88.0699774
P081	202	193	178	132	125	132	0.65346535	0.64766839	0.74157303	86.529563	81.940874	86.529563
P082	207	200	187	151	140	164	0.7294686	0.7	0.87700535	84.6263736	78.4615385	91.9120879
P083	210	200	189	119	135	171	0.56666667	0.675	0.9047619	71.4	81	102.6
P086	212	202	191	153	140	162	0.72169811	0.69306931	0.84816754	85.7472527	78.4615385	90.7912088
P087												
P088	194	185	173	128	118	152	0.65979381	0.63783784	0.87861272	82.0100503	75.6030151	97.3869347
P089	195	179	166	95	83	104	0.48717949	0.46368715	0.62650602	85.9042553	75.0531915	94.0425532
P091	193	184	171	86	91	123	0.44559585	0.49456522	0.71929825	73.1	77.35	104.55
P094	198	188	175	123	163	176	0.62121212	0.86702128	1.00571429	67.8896104	89.9675325	97.1428571
P095												
P096	190	180	167	110	101	132	0.57894737	0.56111111	0.79041916	81.7784257	75.0874636	98.1341108
P097	190	176	149	96	82	110	0.50526316	0.46590909	0.73825503	85	72.6041667	97.3958333
P098	179	168	150	101	121	138	0.56424581	0.7202381	0.92	71.5416667	85.7083333	97.75
P099	201	191	178	134	125	163	0.66666667	0.65445026	0.91573034	80.971564	75.5331754	98.4952607
P100	175	165	154	94	113	107	0.53714286	0.68484848	0.69480519	76.3375796	91.7675159	86.8949045

P101	196	187	176	145	193	180	0.73979592	1.03208556	1.02272727	71.3803089	95.0096525	88.6100386
P102	193	183	172	124	165	131	0.64248705	0.90163934	0.76162791	75.2857143	100.178571	79.5357143
P103	185	172	157	98	129	120	0.52972973	0.75	0.76433121	72.0172911	94.7982709	88.184438
P104												
P105	191	180	168	87	141	143	0.45549738	0.78333333	0.85119048	59.7978437	96.9137466	98.2884097
P114	185	173	161	151	166	194	0.81621622	0.95953757	1.20496894	75.3522505	82.8375734	96.8101761
P116												
P119												
P121	194	183	171	145	175	200	0.74742268	0.95628415	1.16959064	71.1057692	85.8173077	98.0769231
P122												
P123												
P124	179	170	157	78	100	98	0.43575419	0.58823529	0.62420382	72.0652174	92.3913043	90.5434783
P125	179	171	157	79	101	101	0.44134078	0.59064327	0.6433121	71.6903915	91.6548043	91.6548043
P127	178	167	155	132	177	182	0.74157303	1.05988024	1.17419355	68.5539715	91.9246436	94.5213849
P128	180	172	158	135	135	170	0.75	0.78488372	1.07594937	78.2386364	78.2386364	98.5227273
P129	197	187	175	135	148	130	0.68527919	0.79144385	0.74285714	83.3535109	91.3801453	80.2663438
P130	199	187	175	98	135	151	0.49246231	0.72192513	0.86285714	65.078125	89.6484375	100.273438
P134	182	171	158	123	142	165	0.67582418	0.83040936	1.0443038	72.9418605	84.2093023	97.8488372
P135	194	184	173	134	165	170	0.69072165	0.89673913	0.98265896	72.8571429	89.7121535	92.4307036
P136	195	186	174	136	160	161	0.6974359	0.86021505	0.92528736	75.8862144	89.2778993	89.8358862
P137	201	192	179	132	134	153	0.65671642	0.69791667	0.8547486	80.3341289	81.5513126	93.1145585
P138	199	188	176	121	160	160	0.6080402	0.85106383	0.90909091	69.9659864	92.5170068	92.5170068
P139	210	200	186	143	170	173	0.68095238	0.85	0.93010753	75.0308642	89.1975309	90.7716049
P140	200	189	177	142	177	177	0.71	0.93650794	1	73.0040323	90.9979839	90.9979839
P141												
P143	214	207	198	128	166	175	0.59813084	0.80193237	0.88383838	69.5948827	90.2558635	95.1492537
P144	199	188	175	126	167	185	0.63316583	0.88829787	1.05714286	67.2175732	89.0899582	98.6924686
P145	203	196	184	112	121	162	0.55172414	0.61734694	0.88043478	72.3037975	78.1139241	104.582278
P147	205	194	182	132	161	161	0.64390244	0.82989691	0.88461538	74.1409692	90.4295154	90.4295154
P149	209	203	194	145	144	184	0.6937799	0.70935961	0.94845361	78.1712474	77.6321353	99.1966173
P152	207	201	191	144	165	164	0.69565217	0.82089552	0.85863874	77.6321353	88.9534884	88.4143763

P154	180	169	156	136	133	164	0.75555556	0.78698225	1.05128205	80.0923788	78.3256351	96.5819861
P155												
P026	173	166	154	87	119	136	0.50289017	0.71686747	0.88311688	64.8684211	88.7280702	101.403509
P025	205	199	187	89	117	116	0.43414634	0.5879397	0.62032086	70.4813665	92.6552795	91.863354
P159	181	173	163	124	148	136	0.68508287	0.85549133	0.83435583	77.5	92.5	85
P107	185	174	161	112	133	133	0.60540541	0.76436782	0.82608696	75.5555556	89.7222222	89.7222222
P157	172	163	148	122	143	155	0.70930233	0.87730061	1.0472973	74.0714286	86.8214286	94.1071429
P005	175	168	155	105	106	157	0.6	0.63095238	1.01290323	72.7581522	73.451087	108.790761
P004	180	170	156	94	98	124	0.52222222	0.57647059	0.79487179	75.8544304	79.0822785	100.063291
P166	181	171	156	108	132	151	0.59668508	0.77192982	0.96794872	70.4347826	86.0869565	98.4782609
P167												
P027	177	169	156	169	156	174	0.95480226	0.92307692	1.11538462	86.3627255	79.7194389	88.9178357
P148	210	204	192	124	114	155	0.59047619	0.55882353	0.80729167	80.4580153	73.9694656	100.572519
P158	203	195	183	134	141	128	0.66009852	0.72307692	0.69945355	84.7890819	89.2183623	80.9925558

Table 1.

ID #		species	county	locality	loc. #	lat	long	collectors	ridge
P 001	F	<i>P. youngi</i>	Marion	Ocala Nat. For., W of Lake Kerr	4	29.34821	-81.81608	Daniel Dye	Mt.Dora
P 002	M	<i>P. youngi</i>	Marion	Ocala Nat. For., W of Lake Kerr	4	29.34821	-81.81608	Daniel Dye	Mt. Dora
P 003	F	<i>P. profundus</i>	Hernando	Sunshine Grove Rd at Sandy Dr	2	28.5808	-82.4998	Mansell	Southern Brooksville (S)
P 004		<i>P. profundus</i>	Citrus	N Parkwood Ave, S of W Oakhill St.	1	29.01604	-82.53441	Moler	Southern Brooksville (N)
P 005		<i>P. profundus</i>	Citrus	N Parkwood Ave, S of W Oakhill St.	1	29.01604	-82.53441	Moler	Southern Brooksville (N)
P 006		<i>P. profundus</i>	Citrus	N Camae Point	3	29.00785	-82.5243	Moler	Southern Brooksville (N)
P 007		<i>P. youngi</i>	Marion	Ocala Nat. For., W of Lake Kerr	4	29.34821	-81.81608	Daniel Dye	Mt. Dora
P 008	M	<i>P. youngi</i>	Marion	Ocala Nat. For FL Hwy 40, 0.5 mi W FR 11	12	29.1799	-81.7755	Moler	Mt. Dora
P 009	M	<i>P. youngi</i>	Marion	Ocala Nat. For FL Hwy 40, 0.5 mi W FR 11	12	29.1799	-81.7755	Moler	Mt. Dora
P 010	M	<i>P. youngi</i>	Marion	Ocala Nat. For. FR 11, 0.4 mi S Co Rd 314	11	29.3213	-81.7736	Moler	Mt. Dora
P 011	M	<i>P. youngi</i>	Marion	Ocala Nat. For. FR 11, 0.4 mi S Co Rd 314	11	29.3213	-81.7736	Moler	Mt. Dora
P 012	M	<i>P. youngi</i>	Marion	Ocala Nat. For. FR 11, 0.4 mi S Co Rd 314	11	29.3213	-81.7736	Moler	Mt. Dora
P 013	F	<i>P. youngi</i>	Marion	Ocala Nat. For. FR 11, 0.4 mi S Co Rd 314	11	29.3213	-81.7736	Moler	Mt. Dora
P 014	M	<i>P. youngi</i>	Marion	Ocala Nat. For. FL Hwy 40, 0.5 mi W FR 11	12	29.1799	-81.7755	Moler	Mt. Dora
P 015	M	<i>P. youngi</i>	Putnam	Ocala Nat. For. FR 66, 0.3 mi E SR 19	10	29.4342	-81.7325	Moler	Mt. Dora
P 016	M	<i>P. youngi</i>	Putnam	Ocala Nat. For. FR 66, 0.3 mi E SR 19	10	29.4342	-81.7325	Moler	Mt. Dora
P 017	M	<i>P. youngi?</i>	Putnam	Ocala Nat. For. FR 66, 0.3 mi E SR 19	10	29.4342	-81.7325	Moler	Mt. Dora
P 018	M	<i>P. youngi</i>	Putnam	Ocala Nat. For. FR 66, 0.3 mi E SR 19	10	29.4342	-81.7325	Moler	Mt. Dora
P 019	M	<i>P. youngi</i>	Putnam	Ocala Nat. For. FR 66, 0.3 mi E SR 19	10	29.4342	-81.7325	Moler	Mt. Dora

P 020	M	<i>P. youngi?</i>	Putnam	Ocala Nat. For. FR 66, 0.3 mi E SR 19	10	29.4342	-81.7325	Moler	Mt. Dora
P 021	M	<i>P. youngi</i>	Putnam	Ocala Nat. For. FR 66, 0.3 mi E SR 19	10	29.4342	-81.7325	Moler	Mt. Dora
P 022	M	<i>P. profundus</i>	Levy	SR 24, 0.3 mi W Alachua Co line	13	29.5097	-82.5616	Moler	Northern Brooksville
P 023	M	<i>P. youngi?</i>	Putnam	Lake Galilee Dr., 0.15 N FL Hwy 20	9	29.6264	-81.9514	Moler	Trail
P 024	M	<i>P. youngi?</i>	Putnam	Lake Galilee Dr., 0.15 N FL Hwy 20	9	29.6264	-81.9514	Moler	Trail
P 025		<i>P. profundus?</i>	Marion	Hwy 25	7	29.04203	-81.97864	Moler	
P 026		<i>P. profundus</i>	Marion	Indian Lake State Forest	5	29.28685	-82.0661	Moler	
P 027		<i>P. profundus</i>	Marion	Indian Lake State Forest	5	29.28685	-82.0661	Moler	
P 028		<i>P. profundus?</i>	Marion	Silver Spring Shores, Emerald Dr.	6	29.09329	-82.00845	Moler	
P 029		<i>P. profundus?</i>	Marion	Silver Spring Shores, Emerald Dr.	6	29.09329	-82.00845	Moler	
P 030		<i>P. profundus?</i>	Marion	Silver Spring Shores, Emerald Dr.	6	29.09329	-82.00845	Moler	
P 031	M	<i>P. youngi</i>	Putnam	Etoniah State Forest	14	29.7232	-81.8276	Moler	Mt. Dora
P 032	M	<i>P. youngi</i>	Putnam	Etoniah State Forest	14	29.7232	-81.8276	Moler	Mt. Dora
P 033	F	<i>P. profundus</i>	Levy	6.6 mi SW Williston; E Levy St & SE141Cr	16	29.3313	-82.5361	Moler	Northern Brooksville
P 034	M	<i>P. profundus</i>	Levy	4.5 mi WSW Williston; NE 20 St, 0.6 mi W SR121	15	29.3597	-82.5139	Moler	Northern Brooksville
P 035	F	<i>P. profundus</i>	Levy	SR 24, 0.3 mi W Alachua Co line	13	29.5097	-82.5616	Moler	Northern Brooksville
P 036		<i>P. profundus</i>	Levy	SR 24, 0.3 mi W Alachua Co line	13	29.5097	-82.5616	Moler	Northern Brooksville
P 037		<i>P. profundus</i>	Levy	SR 24, 0.3 mi W Alachua Co line	13	29.5097	-82.5616	Moler	Northern Brooksville
P 038		<i>P. profundus</i>	Levy	SR 24, 0.3 mi W Alachua Co line	13	29.5097	-82.5616	Moler	Northern Brooksville
P 039		<i>P. profundus</i>	Levy	SR 24, 0.3 mi W Alachua Co line	13	29.5097	-82.5616	Moler	Northern Brooksville
P 040	M	<i>P. youngi</i>	Marion	SR 42, 2 mi E SE 182 Ave Rd	19	28.9791	-81.7704	Moler	Mt. Dora
P 041		<i>P. youngi?</i>	Lake	SR 44 jct SR 46A	17	28.8509	-81.5303	Moler	Mt. Dora?

P 042		<i>P. youngi?</i>	Lake	W Huff Rd, 2 mi N SR 44	18	28.855	-81.5267	Moler	Mt. Dora?
P 043		<i>P. youngi?</i>	Lake	W Huff Rd, 2 mi N SR 44	18	28.855	-81.5267	Moler	Mt. Dora?
P 044		<i>P. youngi?</i>	Lake	W Huff Rd, 2 mi N SR 44	18	28.855	-81.5267	Moler	Mt. Dora?
P 045		<i>P. youngi</i>	Putnam	Ocala Nat. For.(vial 1)	20	29.4906	-81.8102	Daniel Dye	Mt. Dora
P 046		<i>P. youngi</i>	Marion	Ocala Nat. For. (vial 2)	21	29.4879	-81.8606	Daniel Dye	Mt. Dora
P 047		<i>P. youngi</i>	Putnam	Ocala Nat. For. (vial 4)	23	29.4041	-81.7413	Daniel Dye	Mt. Dora
P 048	F	<i>P. youngi?</i>	Alachua	Holden Park Rd., 0.2 mi NW Putnam Co line	24	29.546	-82.058	Moler	Mt. Dora
P 049	M	<i>P. youngi?</i>	Alachua	Holden Park Rd., 0.2 mi NW Putnam Co line	24	29.546	-82.058	Moler	Mt. Dora
P 050	M	<i>P. youngi</i>	Putnam	Baden Powell Rd at SR 20	25	29.6154	-81.978	Moler	Mt. Dora
P 051	M	<i>P. youngi</i>	Putnam	Baden Powell Rd at SR 20	25	29.6154	-81.978	Moler	Mt. Dora
P 052		<i>P. youngi</i>	Clay	SR 21, 2.2 mi NE SR100	26	29.8099	-82.0091	Moler	Trail
P 053	M	<i>P. youngi</i>	Clay	SR 21, 2.2 mi NE SR100	26	29.8099	-82.0091	Moler	Trail
P 054	M	<i>P. youngi</i>	Clay	SR 21, 2.2 mi NE SR100	26	29.8099	-82.0091	Moler	Trail
P 055	M	<i>P. youngi</i>	Clay	SR 21, 2.2 mi NE SR100	26	29.8099	-82.0091	Moler	Trail
P 056	M	<i>P. profundus</i>	Dixie	US 19, 5.3 mi W Old Town	32	29.60855	-83.07015	Moler	
P 057	F	<i>P. profundus</i>	Dixie	Co Rd 361, 2.1 mi S Co Rd 358 (S. Jena)	33	29.63539	-83.35489	Moler	
P 058	M	<i>P. profundus</i>	Dixie	Co Rd 361, 2.1 mi S Co Rd 358 (S. Jena)	33	29.63539	-83.35489	Moler	
P 059	M	<i>P. profundus</i>	Dixie	Co Rd 361, 2.1 mi S Co Rd 358 (S. Jena)	33	29.63539	-83.35489	Moler	
P 060	M	<i>P. profundus</i>	Dixie	Co Rd 361, 2.1 mi S Co Rd 358 (S. Jena)	33	29.63539	-83.35489	Moler	
P 061	M	<i>P. profundus?</i>	Alachua	Holden Park Rd, 0.25 mi NW Putnam Co line	24	29.5462	-82.058	Moler	Mt. Dora
P 062	M	<i>P. p. dubius</i>	Volusia	De Leon Springs	30	29.1363	-81.3565	Moler	Deland
P 063	F	<i>P. p. dubius</i>	Volusia	De Leon Springs	30	29.1363	-81.3565	Moler	Deland
P 064	M	<i>P. profundus</i>	Gilchrist	SR 26, 3.0 mi W Alachua Co line	34	29.6314	-82.7048	Moler	Northern Brooksville
P 065	F	<i>P. profundus</i>	Gilchrist	SR 26, 3.0 mi W Alachua Co line	34	29.6314	-82.7048	Moler	Northern Brooksville
P 066	M	<i>P. profundus?</i>	Clay	Hwy 21 at Gas Line Rd		29.80734	-82.01186	Moler	Trail

P 067	F	<i>P. profundus</i>	Madison	CR 53, 0.25 mi N Lafayette Co line	38	30.2635	-83.2833	Moler	
P 068	F	<i>P. profundus</i>	Lafayette	CR 53, 1.2 mi N US 27	37	30.1462	-83.2932	Moler	
P 069	F	<i>P. profundus</i>	Dixie	CR 349, 0.15 mi SCR 351A (NE 512 Ave)	43	29.7064	-82.9846	Moler	
P 070	F	<i>P. profundus</i>	Dixie	US 19, 5.3 mi W Old Town		29.60855	-83.07015	Moler	
P 071	M	<i>P. profundus</i>	Taylor	Beach Rd. (CR361) 8.9 mi N Steinhatchee	40	29.7607	-83.4975	Moler	
P 072	F	<i>P. profundus</i>	Taylor	Hampton Spring Rd, 0.4 mi NE Coney Island	42	30.0409	-83.7266	Moler	
P 073	F	<i>P. profundus</i>	Taylor	Hampton Spring Rd, 0.4 mi NE Coney Island	42	30.0409	-83.7266	Moler	
P 074	M	<i>P. profundus</i>	Taylor	FL Hwy 51, 4.9 mi S US 19	39	29.7186	-83.3607	Moler	
P 075	M	<i>P. profundus</i>	Taylor	FL Hwy 51, 4.9 mi S US 19	39	29.7186	-83.3607	Moler	
P 076	F	<i>P. profundus</i>	Gilchrist	CR 340, 2.15 mi E US 129	43	29.7955	-82.8311	Moler	
P 077	F	<i>P. profundus</i>	Gilchrist	CR 340, 2.15 mi E US 129	43	29.7955	-82.8311	Moler	
P 078	M	<i>P. profundus</i>	Hamilton	SR 6, ~1 mi W I-75		30.50637	-83.06778		
P 079	M	<i>P. profundus</i>	Hamilton	CR 141, ~4.8 mi S SR6		30.42931	-83.17017		
P 080	M	<i>P. profundus</i>	Hamilton	CR 141. ~4.8 mi S SR6		30.42931	-81.17017		
P 081	M	<i>P. profundus</i>	Marion	CR 326, ENE 58th Ave.		29.23983	-82.05167		
P 082	M	<i>P. profundus</i>	Hamilton	CR 141, ~ 4.8 mi S SR6		30.42931	-83.17017		
P 083	M	<i>P. profundus</i>	Marion	CR 326 ENE 58th ave.		29.23983	-82.05167		
P 084		<i>Mycotrupes gaigei</i>	Marion	CR 326 ENE 58th ave.					Cotton Plant
P 085		<i>Geotrupes egeriei</i>	Alachua	Fl. 7818 SW CR 346					
P 086	M	<i>P. profundus</i>	Marion	Fl. Co. Rd. 326 E of NE 58th Ave.		29.2403	-82.051	P. Moler	
P 087	M	<i>P. profundus</i>	Hamilton	Fl. SR 6, 0.08 mi E SW CR 143		30.48575	-83.19907	P. Moler	
P 088	M	<i>P. profundus</i>	Hamilton	Fl. SR 6, 0.08 mi E SW CR 144		30.48575	-83.19907	P. Moler	
P 089	M	<i>P. profundus</i>	Hamilton	Fl. SR 6, 0.08 mi E SW CR 145		30.48575	-83.19907	P. Moler	
P 090	F	<i>P. profundus</i>	Marion	Fl. US 27, 1.3 mi N SR 42.		28.9987	-81.99824	P. Moler	
P 091	M	<i>P. profundus</i>	Marion	Fl. US 41 0.5 mi S CR 464		29.23511	-82.44105	P. Moler	

P 092	F	<i>P. profundus</i>	Marion	Fl. US 41 0.5 mi S CR 464 at Rainbow Springs		29.08914	-82.45154	P. Moler	
P 093	M	<i>P. profundus</i>	Madison	I-10 E-bound rest area		30.36751	-83.24831	P. Moler	
P 094	M	<i>P. profundus</i>	Madison	I-10 E-bound rest area		30.36751	-83.24831	P. Moler	
P 095	M	<i>P. profundus</i>	Madison	I-10 E-bound rest area		30.36751	-83.24831	P. Moler	
P 096	M	<i>P. profundus</i>	Madison	Twin Rivers State Forest		30.59969	-83.26375	P. Moler	
P 097	M	<i>P. profundus</i>	Madison	Twin Rivers State Forest		30.59969	-83.26375	P. Moler	
P 098	M	<i>P. profundus</i>	Madison	Twin Rivers State Forest		30.59969	-83.26375	P. Moler	
P 099	M	<i>P. profundus</i>	Madison	Twin Rivers State Forest		30.59969	-83.26375	P. Moler	
P 100	M	<i>P. youngi</i>	Marion	SE 112th Place		29.06292	-81.90894	P. Moler	
P 101	M	<i>P. youngi</i>	Marion	SE 112th Place		29.06292	-81.90894	P. Moler	
P 102	M	<i>P. youngi</i>	Marion	SE 112th Place		29.06292	-81.90894	P. Moler	
P 103	M	<i>P. youngi</i>	Marion	SE 112th Place		29.06292	-81.90894	P. Moler	
P 104	M	<i>P. youngi</i>	Marion	Ocala Nat. Forest, S border		28.9766	-81.772	P. Moler	
P 105	M	<i>P. youngi</i>	Marion	Ocala Nat. Forest, S border		28.9766	-81.772	P. Moler	
P 106	M	<i>P. p. dubius</i>	Volusia	US 17 0.13 mi S Spring Garden Ranch Rd.		29.13355	-81.35167	P. Moler	S Spring Garden Ranch
P 107		<i>P. p. dubius</i>	Volusia	US 17 0.13 mi S Spring Garden Ranch Rd.		29.13355	-81.35167	P. Moler	S Spring Garden Ranch
P 108		<i>P. p. dubius</i>	Putnam	Lake Cond Dr., 0.25 mi S US 17		29.4885	-81.5867	P. Moler	Crescent City
P 109	F	<i>P. profundus</i>	Marion	Fl. US 27 1.2 mi N jct SR		42.99697	-81.99745	P. Moler	
P 110	F	<i>P. profundus</i>	Marion	Fl. US 27 1.2 mi N jct SR		42.99697	-81.99745	P. Moler	
P 111	F	<i>P. profundus</i>	Osceola	US 192 at Co Rd. 500 A		28.22411	-81.18109	P. Moler	
P 112	F	<i>P. profundus</i>	Osceola	US 192 at Co Rd. 500 A		28.22411	-81.18109	P. Moler	
P 113	F	<i>P. profundus</i>	Osceola	US 192 at Co Rd. 500 A		28.22411	-81.18109	P. Moler	
P 114	M	<i>P. profundus</i>	Polk	Tiger Creek Preserve		27.809	-81.4896	P. Moler	Lake Wales
P 115	F	<i>P. profundus</i>	Polk	Tiger Creek Preserve		27.809	-81.4896	P. Moler	Lake Wales
P 116	M	<i>P. profundus</i>	Indian River	Roseland		27.83766	-80.48475	P. Moler	Atlantic Coastal
P 117	F	<i>P. profundus</i>	Indian River	Roseland		27.83766	-80.48475	P. Moler	Atlantic Coastal

P 118	F	<i>P. profundus</i>	Indian River	Roseland		27.83766	-80.48475	P. Moler	Atlantic Coastal
P 119	F	<i>P. profundus</i>	Indian River	Roseland		27.83766	-80.48475	P. Moler	Atlantic Coastal
P 120	M	<i>P. profundus</i>	Indian River	Roseland		27.83766	-80.48475	P. Moler	Atlantic Coastal
P 121	M	<i>P. profundus</i>	Indian River	Roseland		27.83766	-80.48475	P. Moler	Atlantic Coastal
P 122	M	<i>P. profundus</i>	Indian River	Roseland		27.83766	-80.48475	P. Moler	Atlantic Coastal
P 123	M	<i>P. profundus</i>	Highlands	US 27, S side of Josephine Ck.		27.3719	-81.40122	P. Moler	Lake Wales
P 124	F	<i>P. profundus</i>	Highlands	US 27, S side of Josephine Ck.		27.3719	-81.40122	P. Moler	Lake Wales
P 125	M	<i>P. profundus</i>	Polk	Hwy 630 E of Frostproof		27.75978	-81.46325	P. Moler	Lake Wales
P 126	F	<i>P. profundus</i>	Polk	Hwy 580 E of Haines City		28.12093	-81.55084	P. Moler	Lake Wales
P 127	M	<i>P. profundus</i>	Polk	Hwy 580 E of Haines City		28.12093	-81.55084	P. Moler	Lake Wales
P 128	M	<i>P. profundus</i>	Polk	Hwy 580 E of Haines City		28.12093	-81.55084	P. Moler	Lake Wales
P 129	M	<i>P. profundus</i>	Polk	Hwy 580 E of Haines City		28.12093	-81.55084	P. Moler	Lake Wales
P 130	M	<i>P. profundus</i>	Polk	Hwy 580 E of Haines City		28.12093	-81.55084	P. Moler	Lake Wales
P 131	F	<i>P. profundus</i>	Marion	Fl. US 41 0.5 mi S CR 464 at Rainbow Springs		29.08914	-82.45154	P. Moler	
P 132	F	<i>P. profundus</i>	Marion			29.19319	-82.40241	P. Moler	Cotton Plant
P 133	F	<i>P. profundus</i>	Marion			29.19319	-82.40241	P. Moler	Cotton Plant
P 134	M	<i>P. profundus</i>	Marion			29.19319	-82.40241	P. Moler	Cotton Plant
P 135	M	<i>P. profundus</i>	Lake	SE of SR 44 jct SR 46A	17	28.8509	-81.5303	P. Moler	
P 136	M	<i>P. profundus</i>	Lake	SE of SR 44 jct SR 46A	17	28.8509	-81.5303	P. Moler	
P 137	M	<i>P. profundus</i>	Lake	NE jct SR44 and La Place Ct.		28.8516	-81.5656	P. Moler	
P 138	M	<i>P. profundus</i>	Lake	NE jct SR44 and La Place Ct.		28.8516	-81.5656	P. Moler	
P 139	M	<i>P. profundus</i>	Lake	NE jct SR44 and La Place Ct.		28.8516	-81.5656	P. Moler	
P 140	M	<i>P. profundus</i>	Lake	NE jct SR44 and La Place Ct.		28.8516	-81.5656	P. Moler	
P 141	M	<i>P. profundus</i>	Lake	NE jct SR44 and La Place Ct.		28.8516	-81.5656	P. Moler	

P 142	M	<i>P. youngi</i>	Lake	Powerline Rd. E of SR 19		29.051	-81.6341	P. Moler	
P143	M	<i>P. profundus</i>	Marion	CR 326 ENE 58th Ave.		29.2406	-82.0516	P. Moler	
P144	M	<i>P. profundus</i>	Marion	CR 326 ENE 58th Ave.		29.2406	-82.0516	P. Moler	
P145	M	<i>P. profundus</i>	Marion	CR 326 ENE 58th Ave.		29.2406	-82.0516	P. Moler	
P146	M	<i>P. profundus</i>	Marion	CR 326 ENE 58th Ave.		29.2406	-82.0516	P. Moler	Not Tissued
P146		<i>P. youngi</i>	Marion	Ocala NF SR40, 0.4 Mi W FR 11		29.1783	-81.777	P. Moler	
P147		<i>P. youngi</i>	Marion	Ocala Nat. For. FR 11, 0.4 mi S Co Rd 314		29.3219	-81.7727	P. Moler	
148	F	profundeus	MarioN						
149	M	dubius							
150	F	profunus	Marion						
151	M								
152	M								
153	M								
154	M								
155	M								
156	M								
157	M								
158	M								
159	M								
P148	M	<i>P. profundus</i>	Citrus	W. Basket Way at N Travis Dr.		28.9889	-82.4482	P. Moler	Not tissued
P149	M	<i>P. profundus</i>	Citrus	W. Basket Way at N Travis Dr.		28.9889	-82.4482	P. Moler	
P150	F	<i>P. profundus</i>	Levy	US Hwy 41, 1.4 mi N Marion Co. line		29.2349	-82.441	P. Moler	
P151	F	<i>P. profundus</i>	Marion	FL Hwy 4 at SE 172 Terr.		29.1727	-81.866	P. Moler	
P152	M	<i>P. profundus</i>	Marion	FL Hwy 4 at SE 172 Terr.		29.1727	-81.866	P. Moler	
P153	F	<i>P. profundus</i>	Highlands	US 27, S side of Josephine Ck.		27.3719	-81.40122	P. Moler	Lake Wales

Table 2.

